

<p>Form PTO-1449 <b>INFORMATION DISCLOSURE CITATION IN AN APPLICATION</b> <i>(Use several sheets if necessary)</i> <b>DEC 20 2004</b> <b>PTO/SB/08 (2-92)</b></p>	Docket Number 511582003420		Application Number 10/087,190
	Applicant Pia M. CHALLITA-EID et al.		
	Filing Date February 28, 2002	Group Art Unit 1642	
	Mailing Date December 16, 2004		

**U.S. PATENT DOCUMENTS**

Examiner Initials	Ref. No.	Date	Document No.	Name	Class	Subclass	Filing Date If Appropriate
DB	1.	02/13/2003	2003/0032087	Challita-Eid et al.	—	—	
DB	2.	07/17/2001	6,262,333	Endege et al.	—	—	
DB	3.	05/27/2003	6,569,662	Tang et al.	—	—	

**FOREIGN PATENT DOCUMENTS**

Examiner Initials	Ref. No.	Date	Document No.	Country	Class	Subclass	Translation YES NO
DB	4.	07/08/1999	WO 99/33982	WIPO	—	—	
DB	5.	08/05/1999	WO 99/38972	WIPO	—	—	
DB	6.	12/16/1999	WO 99/64576	WIPO	—	—	
DB	7.	01/11/2001	WO 01/02568	WIPO	—	—	
DB	8.	04/05/2001	WO 01/22920	WIPO	—	—	
DB	9.	07/26/2001	WO 01/53312	WIPO	—	—	
DB	10.	09/13/2001	WO 01/66753	WIPO	—	—	
DB	11.	10/11/2001	WO 01/75067	WIPO	—	—	
DB	12.	10/24/2002	WO 02/083876	WIPO	—	—	
DB	13.	11/28/2002	WO 02/095009	WIPO	—	—	

**OTHER DOCUMENTS**

*(including author, title, Date, Pertinent Pages, Etc.)*

Examiner Initials	Ref. No.	Title
DB	14.	Attwood (2000). <i>Science</i> 290:471-473.
DB	15.	Burgess et al. (1990). <i>J. Cell Biol.</i> 111:2129-2138.
DB	16.	Carninci, P. et al. (1999). "High-Efficiency Full-Length cDNA Cloning" Chapter 2 <i>In cDNA Preparation and Characterization - Methods in Enzymology</i> , Weissman, S.M. ed. Academic Press, Inc. 303:19-44.
DB	17.	Carninci, P. et al. (2000). "Normalization and Subtraction of Cap-Trapper-Selected cDNAs to Prepare Full-Length cDNA Libraries for Rapid Discovery of New Genes," <i>Genome Research</i> 10:1617-1630.

EXAMINER: *Pia M. Challita*

DATE CONSIDERED: *4/20/05*

EXAMINER: Initial if citation considered, whether or not the citation conforms with MPEP 609. Draw a line through the citation if not in conformance and not considered. Include a copy of this form with next communication to applicant.

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Docket Number 511582003420

Application Number 10/087,190

Applicant

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INFORMATION DISCLOSURE CITATION  
IN AN APPLICATION

(Use several sheets if necessary)

DEC 20 2004

18.	DGENE Record Accession No. AAX98924 for WO 99/33982, two pages.
19.	DGENE Record Accession No. AAX99020 for WO 99/33982, two pages.
20.	EMBL: Accession No. AAM40043, created on October 21, 2001, European Search Report Reference No. XP 002297423, two pages
21.	EMBL: Accession No. AYO28916, created on March 30, 2001, European Search Report Reference No. XP002297424, one page
22.	EMBL: Accession No. AAZ80011 created on April 7, 2000, European Search Report Reference No. XP-002297421, one page
23.	EMBL: Accession No. BE513408 created on August 8, 2000, European Search Report Reference No. XP-002297422, two pages
24.	GenBank Accession No. BAB27765 created on April 3, 2004 located at < <a href="http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=protein&amp;val=12847934">http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=protein&amp;val=12847934</a> > last visited on November 28, 2004, three pages.
25.	GenBank Accession No. NP_115493 created on October 27, 2004 located at < <a href="http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=protein&amp;val=14149769">http://www.ncbi.nlm.nih.gov/entrez/viewer.fcgi?db=protein&amp;val=14149769</a> > last visited on November 28, 2004, two pages.
26.	Hubert et al. (1999). <i>Proc. Natl. Acad. Sci. USA</i> 96(25):14523-14528.
27.	Klein et al. (1997). <i>Nat. Med.</i> 3:402.
28.	Lazar et al. (1988). <i>Mol. Cell. Biol.</i> 8:1247-1252.
29.	Lerner, (1982). <i>Nature</i> 299:592-596.
30.	Metzler et al. (1997). <i>Nature Structural Biol.</i> 4:527-531.
31.	Mikayama et al. (1993). <i>PNAS</i> 90:10056-10060.
32.	Ngo et al. (1994). <u>The Protein Folding Problem and Tertiary Structure Prediction</u> Birkhauser: Boston, MA pp. 433, 492-495.
33.	Pinto et al. (1996). <i>Clin. Cancer Res.</i> 2(9):1445-1451.
34.	Reiter et al. (1998). <i>Proc. Natl. Acad. Sci. USA</i> 95:1735.
35.	Shibata, K. et al. (2000). "RIKEN Integrated Sequence Analysis (RISA) System - 384-Format Sequencing Pipeline with 384 Multicapillary Sequencer," <i>Genome Research</i> 10:1757-1771.
36.	Skolnick et al. (2000). <i>Trends in Biotech.</i> 18(1):34-39.
37.	Su et al. (1996). <i>Proc. Natl. Acad. Sci. USA</i> 93:7252.
38.	Supplementary Partial Search Report mailed October 6, 2004, for European Patent Application No. 02723291.7, filed February 28, 2002, 7 pages.
39.	DB The FANTOM Consortium and the RIKEN Genome Exploration Research Group Phase I & II Team (December 5, 2002). "Analysis of the Mouse Transcriptome Based on Functional Annotation of 60,770 Full-Length cDNAs," <i>Nature</i> 420:563-573.

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		Applicant	Pia M. CHALLITA-EID et al.
		Filing Date February 28, 2002	Group Art Unit 1642
		Mailing Date December 16, 2004	

JB	40.	The RIKEN Genome Exploration Research Group Phase II Team and the FANTOM Consortium (February 8, 2001). "Functional Annotation of a Full-Length Mouse cDNA Collection," <i>Nature</i> 409:685-690.
	41.	Tsubouchi, H. et al. (May 2002). "The Mnd1 Protein Forms a Complex with Hop2 to Promote Homologous Chromosome Pairing and Meiotic Double-Strand Break Repair," <i>Molecular and Cellular Biology</i> 22(9):3078-3088.

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<p><i>Paul Blahd</i></p> <p>EXAMINER: Initial if citation considered, whether or not the citation conforms with MPEP 609. Draw a line through the citation if not in conformance and not considered. Include a copy of this form with next communication to applicant.</p>	